

CRF Errors Corrected by the STIC System's Branch

Lincoln Stale 1814 11/14
 305-4507
 CRF Processing Date: 11/12/99
 Edited by: _____
 Verified by: lw (STIC staff)

Serial Number: 08/898,560

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☒ Other: Seq 1, 2 - changed KEY to "NAME/KEY" under (ix) FEATURES

***Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.**

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/898,560DATE: 11/12/97
TIME: 21:24:16

INPUT SET: S21525.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

SEQUENCE LISTING

1
2
3 (1) General Information:
4
5 (i) APPLICANT: Hiroyuki NAKANE, Chikara OHTO, Shinichi OHNUMA,
6 Kazutake HIROOKA, Tokuzo NISHINO
7
8 (ii) TITLE OF INVENTION: Farnesyl Diphosphate Synthase
9
10 (iii) NUMBER OF SEQUENCES: 14
11
12 (iv) CORRESPONDENCE ADDRESS:
13 (A) ADDRESSEE: Kenyon & Kenyon
14 (B) STREET: One Broadway
15 (C) CITY: New York
16 (D) STATE: NY
17 (E) COUNTRY: USA
18 (F) ZIP: 10004
19
20 (v) COMPUTER READABLE FORM:
21 (A) MEDIUM TYPE: 3" Floppy disk
22 (B) COMPUTER: IBM PC compatible
23 (C) OPERATING SYSTEM: PC-DOS/MS-DOS 6.2
24 (D) SOFTWARE: WordPerfect 6.1 Windows
25
26 (vi) CURRENT APPLICATION DATA:
27 (A) APPLICATION NUMBER: Not Yet Issued
28 (B) FILING DATE: Concurrent Herewith
29 (C) CLASSIFICATION: Not Yet Issued
30
31 (vii) PRIOR APPLICATION DATA:
32 (A) APPLICATION NUMBER: JP 8-213211
33 (B) FILING DATE: 24-JUL-96
34
35 (viii) ATTORNEY/AGENT INFORMATION:
36 (A) NAME: Edward W. Greason
37 (B) REGISTRATION NUMBER: 18,918
38 (C) REFERENCE/DOCKET NUMBER: 77670/495
39
40 (ix) TELECOMMUNICATION INFORMATION:
41 (A) TELEPHONE: (212)425-7200
42 (B) TELEFAX: (212)425-5288
43
44 (2) INFORMATION FOR SEQ ID NO:1:
45
46 (i) SEQUENCE CHARACTERISTICS:

RAW SEQUENCE LISTING PATENT APPLICATION US/08/898,560

DATE: 11/12/97
TIME: 21:24:19

INPUT SET: S21525.raw

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47      (A) LENGTH: 330 amino acids
48      (B) TYPE: amino acid
49      (D) TOPOLOGY: linear
50
51      (ii) MOLECULE TYPE: protein
52
53      (vi) ORIGINAL SOURCE:
54          (A) ORGANISM: Sulfolobus acidocaldarius
55          (B) STRAIN: ATCC 33909
56
57      (ix) FEATURE:
58          (A) NAME/KEY: Asp-rich domain
59          (B) LOCATION: 82-86
60
61      (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
62
63      Met Ser Tyr Phe Asp Asn Tyr Phe Asn Glu Ile Val Asn Ser Val Asn
64              5              10              15
65      Asp Ile Ile Lys Ser Tyr Ile Ser Gly Asp Val Pro Lys Leu Tyr Glu
66              20              25              30
67      Ala Ser Tyr His Leu Phe Thr Ser Gly Gly Lys Arg Leu Arg Pro Leu
68              35              40              45
69      Ile Leu Thr Ile Ser Ser Asp Leu Phe Gly Gly Gln Arg Glu Arg Ala
70              50              55              60
71      Tyr Tyr Ala Gly Ala Ala Ile Glu Val Leu His Thr Phe Thr Leu Val
72              65              70              75              80
73      His Asp Asp Ile Met Asp Gln Asp Asn Ile Arg Arg Gly Leu Pro Thr
74              85              90              95
75      Val His Val Lys Tyr Gly Leu Pro Leu Ala Ile Leu Ala Gly Asp Leu
76              100             105             110
77      Leu His Ala Lys Ala Phe Gln Leu Leu Thr Gln Ala Leu Arg Gly Leu
78              115             120             125
79      Pro Ser Glu Thr Ile Ile Lys Ala Phe Asp Ile Phe Thr Arg Ser Ile
80              130             135             140
81      Ile Ile Ile Ser Glu Gly Gln Ala Val Asp Met Glu Phe Glu Asp Arg
82      145             150             155             160
83      Ile Asp Ile Lys Glu Gln Glu Tyr Leu Asp Met Ile Ser Arg Lys Thr
84              165             170             175
85      Ala Ala Leu Phe Ser Ala Ser Ser Ser Ile Gly Ala Leu Ile Ala Gly
86              180             185             190
87      Ala Asn Asp Asn Asp Val Arg Leu Met Ser Asp Phe Gly Thr Asn Leu
88              195             200             205
89      Gly Ile Ala Phe Gln Ile Val Asp Asp Ile Leu Gly Leu Thr Ala Asp
90              210             215             220
91      Glu Lys Glu Leu Gly Lys Pro Val Phe Ser Asp Ile Arg Glu Gly Lys
92      225             230             235             240
93
94      Lys Thr Ile Leu Val Ile Lys Thr Leu Glu Leu Cys Lys Glu Asp Glu
95              245             250             255
96      Lys Lys Ile Val Leu Lys Ala Leu Gly Asn Lys Ser Ala Ser Lys Glu
97              260             265             270
98      Glu Leu Met Ser Ser Ala Asp Ile Ile Lys Lys Tyr Ser Leu Asp Tyr
99              275             280             285

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RAW SEQUENCE LISTING
PATENT APPLICATION US/08/898,560DATE: 11/12/97
TIME: 21:24:23

INPUT SET: S21525.raw

100 Ala Tyr Asn Leu Ala Glu Lys Tyr Tyr Lys Asn Ala Ile Asp Ser Leu
101 290 295 300
102 Asn Gln Val Ser Ser Lys Ser Asp Ile Pro Gly Lys Ala Leu Lys Tyr
103 305 310 315 320
104 Leu Ala Glu Phe Thr Ile Arg Arg Arg Lys
105 325 330
106

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 993 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: double

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: genomic DNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Sulfolobus acidocaldarius

(B) STRAIN: ATCC 33909

(ix) FEATURE:

(A) NAME/KEY: Asp-rich domain coding

(B) LOCATION: 246-258

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

127 ATGAGTTACT TTGACAACTA TTTTAATGAG ATTGTTAATT CTGTAAACGA CATTATTAAG 60
128
129 AGCTATATAT CTGGAGATGT TCCTAAACTA TATGAAGCCT CATATCATTT GTTTACATCT 120
130
131 GGAGGTAAGA GGTAAAGACC ATTAATCTTA ACTATATCAT CAGATTTATT CGGAGGACAG 180
132
133 AGAGAAAGAG CTTATTATGC AGGTGCAGCT ATTGAAGTTC TTCATACTTT TACGCTTGTG 240
134
135 CATGATGATA TTATGGATCA AGATAATATC AGAAGAGGGT TACCCACAGT CCACGTGAAA 300
136
137 TACGGCTTAC CCTTAGCAAT ATTAGCTGGG GATTTACTAC ATGCAAAGGC TTTTCAGCTC 360
138
139 TTAACCCAGG CTCTTAGAGG TTTGCCAAGT GAAACCATAA TTAAGGCTTT CGATATTTTC 420
140
141 ACTCGTTCAA TAATAATTAT ATCCGAAGGA CAGGCAGTAG ATATGGAATT TGAGGACAGA 480
142
143 ATTGATATAA AGGAGCAGGA ATACCTTGAC ATGATCTCAC GTAAGACAGC TGCATTATTC 540
144
145 TCGGCATCCT CAAGTATAGG CGCACTTATT GCTGGTGCTA ATGATAATGA TGTAAGACTG 600
146
147 ATGTCTGATT TCGGTACGAA TCTAGGTATT GCATTTTCTA TTGTTGACGA TATCTTAGGT 660
148
149 CTAACAGCAG ACGAAAAGGA ACTTGGAAG CCTGTTTTTA GTGATATTAG GGAGGGTAAA 720
150
151 AAGACTATAC TTGTAATAAA AACACTGGAG CTTTGTAAG AGGACGAGAA GAAGATTGTC 780
152

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/898,560DATE: 11/12/97
TIME: 21:24:26

INPUT SET: S21525.raw

153 CTAAAGGCGT TAGGTAATAA GTCAGCCTCA AAAGAAGAAT TAATGAGCTC AGCAGATATA 840
154
155 ATTAAGAAAT ACTCTTTAGA TTATGCATAC AATTTAGCAG AGAAATATTA TAAAAATGCT 900
156
157 ATAGACTCTT TAAATCAAGT CTCCTCTAAG AGTGATATAC CTGGAAAGGC TTTAAAATAT 960
158
159 CTAGCTGAAT TTACGATAAG AAGGAGAAAA TAA 993
160

161

162 (2) INFORMATION FOR SEQ ID NO:3:

163

164 (i) SEQUENCE CHARACTERISTICS:

165 (A) LENGTH:37

166 (B) TYPE: nucleic acid

167 (C) STRANDEDNESS: single

168 (D) TOPOLOGY: linear

169

170 (ii) MOLECULE TYPE: cDNA

171

172 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

173

174 CATACTTTTT TCCTTGTGGC TGATGATATC ATGGATC 37

175

176

177 (2) INFORMATION FOR SEQ ID NO:4:

178

179 (i) SEQUENCE CHARACTERISTICS:

180 (A) LENGTH:37

181 (B) TYPE: nucleic acid

182 (C) STRANDEDNESS: single

183 (D) TOPOLOGY: linear

184

185 (ii) MOLECULE TYPE: cDNA

186

187 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

188

189 CATACTTTTT TCCTTGTGCT TGATGATATC ATGGATC 37

190

191 (2) INFORMATION FOR SEQ ID NO:5:

192

193 (i) SEQUENCE CHARACTERISTICS:

194 (A) LENGTH:37

195 (B) TYPE: nucleic acid

196 (C) STRANDEDNESS: single

197 (D) TOPOLOGY: linear

198

199 (ii) MOLECULE TYPE: cDNA

200

201 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

202

203 CATACTTATT TCCTTGTGCT TGATGATATC ATGGATC 37

204

205

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/898,560DATE: 11/12/97
TIME: 21:24:29

INPUT SET: S21525.raw

206 (2) INFORMATION FOR SEQ ID NO:6:

207

208 (i) SEQUENCE CHARACTERISTICS:

209 (A) LENGTH:37

210 (B) TYPE: nucleic acid

211 (C) STRANDEDNESS: single

212 (D) TOPOLOGY: linear

213

214 (ii) MOLECULE TYPE: cDNA

215

216 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

217

218 CATACTTATT TCCTTGTGGC TGATGATATC ATGGATC

37

219

220

221 (2) INFORMATION FOR SEQ ID NO:7:

222

223 (i) SEQUENCE CHARACTERISTICS:

224 (A) LENGTH: 36

225 (B) TYPE: nucleic acid

226 (C) STRANDEDNESS: single

227 (D) TOPOLOGY: linear

228

229 (ii) MOLECULE TYPE: cDNA

230

231

232 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

233

234 GTTCTTCATA CTTATTCGCT TATTCATGAT AGTATT

36

235

236

237

238

239 (2) INFORMATION FOR SEQ ID NO:8:

240

241 (i) SEQUENCE CHARACTERISTICS:

242 (A) LENGTH: 33

243 (B) TYPE: nucleic acid

244 (C) STRANDEDNESS: single

245 (D) TOPOLOGY: linear

246

247 (ii) MOLECULE TYPE: cDNA

248

249 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

250

251 ATTCATGATG ATCTTCCATC GATGGATCAA GAT

33

252

253

254 (2) INFORMATION FOR SEQ ID NO:9:

255

256 (i) SEQUENCE CHARACTERISTICS:

257 (A) LENGTH: 27

258 (B) TYPE: nucleic acid

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/898,560DATE: 11/12/97
TIME: 21:24:33

INPUT SET: S21525.raw

***** PREVIOUSLY ERRORED SEQUENCES - EDITED *****

44 (2) INFORMATION FOR SEQ ID NO:1:

45

46 (i) SEQUENCE CHARACTERISTICS:

47 (A) LENGTH: 330 amino acids

48 (B) TYPE: amino acid

49 (D) TOPOLOGY: linear

50

51 (ii) MOLECULE TYPE: protein

52

53 (vi) ORIGINAL SOURCE:

54 (A) ORGANISM: Sulfolobus acidocaldarius

55 (B) STRAIN: ATCC 33909

56

57 (ix) FEATURE:

58 (A) NAME/KEY: Asp-rich domain

59 (B) LOCATION: 82-86

60

61 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

62

63 Met Ser Tyr Phe Asp Asn Tyr Phe Asn Glu Ile Val Asn Ser Val Asn

64 5 10 15

65 Asp Ile Ile Lys Ser Tyr Ile Ser Gly Asp Val Pro Lys Leu Tyr Glu

66 20 25 30

67 Ala Ser Tyr His Leu Phe Thr Ser Gly Gly Lys Arg Leu Arg Pro Leu

68 35 40 45

69 Ile Leu Thr Ile Ser Ser Asp Leu Phe Gly Gly Gln Arg Glu Arg Ala

70 50 55 60

71 Tyr Tyr Ala Gly Ala Ala Ile Glu Val Leu His Thr Phe Thr Leu Val

72 65 70 75 80

73 His Asp Asp Ile Met Asp Gln Asp Asn Ile Arg Arg Gly Leu Pro Thr

74 85 90 95

75 Val His Val Lys Tyr Gly Leu Pro Leu Ala Ile Leu Ala Gly Asp Leu

76 100 105 110

77 Leu His Ala Lys Ala Phe Gln Leu Leu Thr Gln Ala Leu Arg Gly Leu

78 115 120 125

79 Pro Ser Glu Thr Ile Ile Lys Ala Phe Asp Ile Phe Thr Arg Ser Ile

80 130 135 140

81 Ile Ile Ile Ser Glu Gly Gln Ala Val Asp Met Glu Phe Glu Asp Arg

82 145 150 155 160

83 Ile Asp Ile Lys Glu Gln Glu Tyr Leu Asp Met Ile Ser Arg Lys Thr

84 165 170 175

85 Ala Ala Leu Phe Ser Ala Ser Ser Ser Ile Gly Ala Leu Ile Ala Gly

86 180 185 190

87 Ala Asn Asp Asn Asp Val Arg Leu Met Ser Asp Phe Gly Thr Asn Leu

88 195 200 205

89 Gly Ile Ala Phe Gln Ile Val Asp Asp Ile Leu Gly Leu Thr Ala Asp

90 210 215 220

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/898,560DATE: 11/12/97
TIME: 21:24:36

INPUT SET: S21525.raw

91 Glu Lys Glu Leu Gly Lys Pro Val Phe Ser Asp Ile Arg Glu Gly Lys
92 225 230 235 240
93
94 Lys Thr Ile Leu Val Ile Lys Thr Leu Glu Leu Cys Lys Glu Asp Glu
95 245 250 255
96 Lys Lys Ile Val Leu Lys Ala Leu Gly Asn Lys Ser Ala Ser Lys Glu
97 260 265 270
98 Glu Leu Met Ser Ser Ala Asp Ile Ile Lys Lys Tyr Ser Leu Asp Tyr
99 275 280 285
100 Ala Tyr Asn Leu Ala Glu Lys Tyr Tyr Lys Asn Ala Ile Asp Ser Leu
101 290 295 300
102 Asn Gln Val Ser Ser Lys Ser Asp Ile Pro Gly Lys Ala Leu Lys Tyr
103 305 310 315 320
104 Leu Ala Glu Phe Thr Ile Arg Arg Arg Lys
105 325 330
106

107 (2) INFORMATION FOR SEQ ID NO:2:
108

109 (i) SEQUENCE CHARACTERISTICS:

110 (A) LENGTH: 993 base pairs

111 (B) TYPE: nucleic acid

112 (C) STRANDEDNESS: double

113 (D) TOPOLOGY: linear
114115 (ii) MOLECULE TYPE: genomic DNA
116

117 (vi) ORIGINAL SOURCE:

118 (A) ORGANISM: Sulfolobus acidocaldarius

119 (B) STRAIN: ATCC 33909
120

121 (ix) FEATURE:

122 (A) NAME/KEY: Asp-rich domain coding

123 (B) LOCATION: 246-258
124125 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
126127 ATGAGTTACT TTGACAACTA TTTTAATGAG ATTGTTAATT CTGTAAACGA CATTATTAAG 60
128129 AGCTATATAT CTGGAGATGT TCCTAAACTA TATGAAGCCT CATATCATTT GTTTACATCT 120
130131 GGAGGTAAGA GGTAAAGACC ATTAATCTTA ACTATATCAT CAGATTTATT CGGAGGACAG 180
132133 AGAGAAAGAG CTTATTATGC AGGTGCAGCT ATTGAAGTTC TTCATACTTT TACGCTTGTG 240
134135 CATGATGATA TTATGGATCA AGATAATATC AGAAGAGGGT TACCCACAGT CCACGTGAAA 300
136137 TACGGCTTAC CCTTAGCAAT ATTAGCTGGG GATTTACTAC ATGCAAAGGC TTTTCAGCTC 360
138139 TTAACCCAGG CTCTTAGAGG TTTGCCAAGT GAAACCATAA TTAAGGCTTT CGATATTTTC 420
140141 ACTCGTTCAA TAATAATTAT ATCCGAAGGA CAGGCAGTAG ATATGGAATT TGAGGACAGA 480
142

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/898,560DATE: 11/12/97
TIME: 21:24:40

INPUT SET: S21525.raw

143	ATTGATATAA	AGGAGCAGGA	ATACCTTGAC	ATGATCTCAC	GTAAGACAGC	TGCATTATTC	540
144							
145	TCGGCATCCT	CAAGTATAGG	CGCACTTATT	GCTGGTGCTA	ATGATAATGA	TGTAAGACTG	600
146							
147	ATGTCTGATT	TCGGTACGAA	TCTAGGTATT	GCATTTTCAGA	TTGTTGACGA	TATCTTAGGT	660
148							
149	CTAACAGCAG	ACGAAAAGGA	ACTTGAAAAG	CCTGTTTTTA	GTGATATTAG	GGAGGGTAAA	720
150							
151	AAGACTATAC	TTGTAATAAA	AACACTGGAG	CTTTGTAAAG	AGGACGAGAA	GAAGATTGTC	780
152							
153	CTAAAGGCGT	TAGGTAATAA	GTCAGCCTCA	AAAGAAGAAT	TAATGAGCTC	AGCAGATATA	840
154							
155	ATTAAGAAAT	ACTCTTTAGA	TTATGCATAC	AATTTAGCAG	AGAAATATTA	TAAAAATGCT	900
156							
157	ATAGACTCTT	TAAATCAAGT	CTCCTCTAAG	AGTGATATAC	CTGGAAAGGC	TTTAAAATAT	960
158							
159	CTAGCTGAAT	TTACGATAAG	AAGGAGAAAA	TAA			993
160							
161							

PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/898,560

DATE: 11/12/97
TIME: 21:24:41

INPUT SET: S21525.raw

Line	Error	Original Text
27	Wrong application Serial Number	(A) APPLICATION NUMBER: Not Yet Issued
29	Wrong Classification	(C) CLASSIFICATION: Not Yet Issued

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/898,560DATE: 11/12/97
TIME: 16:24:22

INPUT SET: S21525.raw

This Raw Listing contains the General
Information Section and those Sequences
containing ERRORS.

SEQUENCE LISTING

Does Not Comply
Corrected Diskette Needed

(1) General Information:

(i) APPLICANTS: Hiroyuki NAKANE, Chikara OHTO, Shinichi OHNUMA,
Kazutake HIROOKA, Tokuzo NISHINO

(ii) TITLE OF INVENTION: Farnesyl Diphosphate Synthase

(iii) NUMBER OF SEQUENCES: 14

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Kenyon & Kenyon

(B) STREET: One Broadway

(C) CITY: New York

(D) STATE: NY

(E) COUNTRY: USA

(F) ZIP: 10004

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: 3" Floppy disk

(B) COMPUTER: IBM PC compatible

(C) OPERATING SYSTEM: PC-DOS/MS-DOS 6.2

(D) SOFTWARE: WordPerfect 6.1 Windows

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: Not Yet Issued

(B) FILING DATE: Concurrent Herewith

(C) CLASSIFICATION: Not Yet Issued

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: JP 8-213211

(B) FILING DATE: 24-JUL-96

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Edward W. Greason

(B) REGISTRATION NUMBER: 18,918

(C) REFERENCE/DOCKET NUMBER: 77670/495

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: (212)425-7200

(B) TELEFAX: (212)425-5288

ERRORED SEQUENCES FOLLOW:

PAGE 2

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/898,560DATE: 11/12/97
TIME: 16:24:25

INPUT SET: S21525.raw

(2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 330 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Sulfolobus acidocaldarius

(B) STRAIN: ATCC 33909

(ix) FEATURE:

(A) KEY: Asp-rich domain

(B) LOCATION: 82-86

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

Met Ser Tyr Phe Asp Asn Tyr Phe Asn Glu Ile Val Asn Ser Val Asn
5 10 15
Asp Ile Ile Lys Ser Tyr Ile Ser Gly Asp Val Pro Lys Leu Tyr Glu
20 25 30
Ala Ser Tyr His Leu Phe Thr Ser Gly Gly Lys Arg Leu Arg Pro Leu
35 40 45
Ile Leu Thr Ile Ser Ser Asp Leu Phe Gly Gly Gln Arg Glu Arg Ala
50 55 60
Tyr Tyr Ala Gly Ala Ala Ile Glu Val Leu His Thr Phe Thr Leu Val
65 70 75 80
His Asp Asp Ile Met Asp Gln Asp Asn Ile Arg Arg Gly Leu Pro Thr
85 90 95
Val His Val Lys Tyr Gly Leu Pro Leu Ala Ile Leu Ala Gly Asp Leu
100 105 110
Leu His Ala Lys Ala Phe Gln Leu Thr Gln Ala Leu Arg Gly Leu
115 120 125
Pro Ser Glu Thr Ile Ile Lys Ala Phe Asp Ile Phe Thr Arg Ser Ile
130 135 140
Ile Ile Ile Ser Glu Gly Gln Ala Val Asp Met Glu Phe Glu Asp Arg
145 150 155 160
Ile Asp Ile Lys Glu Gln Glu Tyr Leu Asp Met Ile Ser Arg Lys Thr
165 170 175
Ala Ala Leu Phe Ser Ala Ser Ser Ser Ile Gly Ala Leu Ile Ala Gly
180 185 190
Ala Asn Asp Asn Asp Val Arg Leu Met Ser Asp Phe Gly Thr Asn Leu
195 200 205
Gly Ile Ala Phe Gln Ile Val Asp Asp Ile Leu Gly Leu Thr Ala Asp
210 215 220
Glu Lys Glu Leu Gly Lys Pro Val Phe Ser Asp Ile Arg Glu Gly Lys
225 230 235 240
Lys Thr Ile Leu Val Ile Lys Thr Leu Glu Leu Cys Lys Glu Asp Glu
245 250 255
Lys Lys Ile Val Leu Lys Ala Leu Gly Asn Lys Ser Ala Ser Lys Glu

PAGE: 3

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/898,560

DATE: 11/12/97

TIME: 16:24:29

INPUT SET: S21525.raw

97 260 265
98 Glu Leu Met Ser Ser Ala Asp Ile Ile Lys Lys Tyr Ser Leu Asp Tyr
99 275 280 285
100 Ala Tyr Asn Leu Ala Glu Lys Tyr Tyr Lys Asn Ala Ile Asp Ser Leu
101 290 295 300
102 Asn Gln Val Ser Ser Lys Ser Asp Ile Pro Gly Lys Ala Leu Lys Tyr
103 305 310 315 320
104 Leu Ala Glu Phe Thr Ile Arg Arg Arg Lys
105 325 330
106

107 (2) INFORMATION FOR SEQ ID NO:2:

108

109 (i) SEQUENCE CHARACTERISTICS:

110 (A) LENGTH: 993 base pairs

111 (B) TYPE: nucleic acid

112 (C) STRANDEDNESS: double

113 (D) TOPOLOGY: linear

114

115 (ii) MOLECULE TYPE: genomic DNA

116

117 (vi) ORIGINAL SOURCE:

118 (A) ORGANISM: Sulfolobus acidocaldarius

119 (B) STRAIN: ATCC 33909

120

121 (ix) FEATURE:

--> 122 (A) KEY: Asp-rich domain coding

123 (B) LOCATION: 246-258

124

125 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

126

127 ATGAGTTACT TTGACAACTA TTTTAATGAG ATTGTTAATT CTGTAAACGA CATTATTAAG 60

128

129 AGCTATATAT CTGGAGATGT TCCTAAACTA TATGAAGCCT CATATCATTT GTTTACATCT 120

130

131 GGAGGTAAGA GGTAAAGACC ATTAATCTTA ACTATATCAT CAGATTTATT CGGAGGACAG 180

132

133 AGAGAAAGAG CTTATTATGC AGGTGCAGCT ATTGAAGTTC TTCATACTTT TACGCTTGTG 240

134

135 CATGATGATA TTATGGATCA AGATAATATC AGAAGAGGGT TACCCACAGT CCACGTGAAA 300

136

137 TACGGCTTAC CCTTAGCAAT ATTAGCTGGG GATTTACTAC ATGCAAAGGC TTTTCAGCTC 360

138

139 TTAACCCAGG CTCTTAGAGG TTTGCCAAGT GAAACCATAA TTAAGGCTTT CGATATTTTC 420

140

141 ACTCGTTCAA TAATAATTAT ATCCGAAGGA CAGGCAGTAG ATATGGAATT TGAGGACAGA 480

142

143 ATTGATATAA AGGAGCAGGA ATACCTTGAC ATGATCTCAC GTAAGACAGC TGCATTATTC 540

144

145 TCGGCATCCT CAAGTATAGG CGCACTTATT GCTGGTGCTA ATGATAATGA TGTAAGACTG 600

146

147 ATGTCTGATT TCGGTACGAA TCTAGGTATT GCATTCAGA TTGTTGACGA TATCTTAGGT 660

148

PAGE: 4

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/898,560DATE: 11/12/97
TIME: 16:24:32

INPUT SET: S21525.raw

149	CTAACAGCAG	ACGAAAAGGA	ACTTGGAAAG	CCTGTTTTTA	GTGATATTAG	GGAGGGTAAA	720
150							
151	AAGACTATAC	TTGTAATAAA	AACACTGGAG	CTTTGTAAAG	AGGACGAGAA	GAAGATTGTC	780
152							
153	CTAAAGGCGT	TAGGTAATAA	GTCAGCCTCA	AAAGAAGAAT	TAATGAGCTC	AGCAGATATA	840
154							
155	ATTAAGAAAT	ACTCTTTAGA	TTATGCATAC	AATTTAGCAG	AGAAATATTA	TAAAAATGCT	900
156							
157	ATAGACTCTT	TAAATCAAGT	CTCCTCTAAG	AGTGATATAC	CTGGAAAGGC	TTTAAAAATAT	960
158							
159	CTAGCTGAAT	TTACGATAAG	AAGGAGAAAA	TAA			993
160							
161							

PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/898,560

DATE: 11/12/97
TIME: 16:24:35

INPUT SET: S21525.raw

Line	Error
27	Wrong application Serial Number
29	Wrong Classification
58	Unknown or Misplaced Identifier
122	Unknown or Misplaced Identifier

Original Text

(A) APPLICATION NUMBER: Not Yet Issued
(C) CLASSIFICATION: Not Yet Issued
(A) KEY: Asp-rich domain
(A) KEY: Asp-rich domain coding